



PCT

**RAW SEQUENCE LISTING**  
**PATENT APPLICATION: US/10/500,586**

**DATE: 07/08/2004**  
**TIME: 16:27:07**

**Input Set : A:\280277 Sequence Listings.txt**  
**Output Set: N:\CRF4\07082004\J500586.raw**

1 <110> APPLICANT: KIM, Bum-Joon  
 2 BIOMEDLAB CORPORATION  
 4 <120> TITLE OF INVENTION: PRIMERS FOR AMPLIFYING HSP 65 GENE OF MYCOBACTERIAL SPECIES,  
 HSP  
 5 65 GENE FRAGMENTS AND METHOD OF IDENTIFYING MYCOBACTERIAL SPECIES  
 6 WITH THE SAME  
 8 <130> FILE REFERENCE: OPP021096KR  
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/500,586  
 C--> 10 <141> CURRENT FILING DATE: 2004-06-30  
 10 <150> PRIOR APPLICATION NUMBER: KR 10-2002-0004297  
 11 <151> PRIOR FILING DATE: 2002-01-24  
 13 <150> PRIOR APPLICATION NUMBER: KR 10-2002-0011648  
 14 <151> PRIOR FILING DATE: 2002-03-05  
 16 <160> NUMBER OF SEQ ID NOS: 56  
 18 <170> SOFTWARE: KopatentIn 1.71  
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 21 <211> LENGTH: 604  
 22 <212> TYPE: DNA  
 23 <213> ORGANISM: Mycobacterium abscessus  
 25 <400> SEQUENCE: 1  
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 28 cgacgtcgcg ggtgacggca ccaccacccg caccgtgctc gcccaggctc tggtaagga 120  
 30 aggtctgcgt aacgtcgccg cccgcgccaa cccgcgtcgcc ctgaagcgcg gtatcgagaa 180  
 32 gggcgtcgag aaggtcacccg agacgctgtc gaagagcgcc aaggaggctcg agaccaagga 240  
 34 gcagatcgcg gccacggccg gtatctccgc gggcgaccag tccatcgccg acctgtatcgc 300  
 36 cgaggccatg gacaagggtt gtaacgaggg tgcgtatcacc gtcgaggagt ccaacacctt 360  
 38 cggcctgcag ctggagctca cccgggttat gcgcttcgac aagggttaca tctcggtt 420  
 40 cttcgtgacc gacgcccggc gtcaggaagc cgtcctggag gatcccttaca tctgtgtt 480  
 42 cagctccaag gtgtcgaccg tcaaggatot gtttcgttg ctggagaagg tcattcaggc 540  
 44 cggcaagccg ctgctgatca tcggcgagga cgtcgaggcc gaggctctt ccactcttgtt 600  
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 49 <210> SEQ ID NO: 2  
 50 <211> LENGTH: 604  
 51 <212> TYPE: DNA  
 52 <213> ORGANISM: Mycobacterium africanum  
 54 <400> SEQUENCE: 2  
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 57 tgacgtcgcc ggtgacggca ccacgacggc caccgtgctg gcccaggctg tggtaagga 120  
 59 gggcctgcgc aacgtcgccg cccgcgccaa cccgcgtcggt ctcaaacgcg gcatcgaaaa 180  
 61 gggcgtggag aaggtcacccg agacccctgtc caaggccggc aaggaggctcg agaccaagga 240  
 63 gcagattcgcg gccaccccgccg cgttgcggc gggcgaccag tccatcgccg acctgtatcgc 300  
 65 cgaggccatg gacaagggtt gcaacggccg cgtcatcacc gtcgaggagt ccaacacctt 360  
 67 tggcgtcgcc ctgcgtatca cccgggttat gcggttcgac aagggttaca tctcggtt 420  
 69 cttcgtgacc gacccggcc gtcaggaggg gttccgttg gacccttaca tctgtgtt 480  
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71	cagctccaag	gtgtccactg	tcaaggatct	gctgccgctg	ctcgagaagg	tcatcgagc	540
73	cggtaagccg	ctgctgatca	tcgcccagga	cgtcgagggc	gaggcgctgt	ccaccctggt	600
75	cgtc						604
78	<210>	SEQ ID NO:	3				
79	<211>	LENGTH:	604				
80	<212>	TYPE:	DNA				
81	<213>	ORGANISM:	Mycobacterium asiaticum				
83	<400>	SEQUENCE:	3				
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86	cgacgtggcc	ggtgacggca	ccacgacggc	caccgtgctg	gcacaggcgc	tggtaaagga	120
88	gggcctgcgc	aacgttgcgg	caggcgccaa	cccgcgtggc	ctgaagcgcg	gcatcgagaa	180
90	ggccgtcgag	aaggtcaccc	agaccctgct	cagctcgcc	aaggacgtcg	agaccaagga	240
92	gcagatcgcg	gccaccgggg	gtatttccgc	gggcgaccag	tcgatcgcc	acctgtatgc	300
94	cgaggcgatg	gacaaagtcg	gcaacgaggg	tgtcatcacc	gtcgaggagt	ccaacacctt	360
96	cggcctgcag	ctcgagctca	ccgagggtat	gcccgtcgac	aagggttaca	tctcggttta	420
98	cttcgtcacc	gacgcccggc	gtcaggaagc	cgtcctggag	gacccttaca	tcctgctgg	480
100	ttccagcaag	gtgtcgaccg	tcaaggacct	gctgccgctg	ctggagaagg	tcatccaggc	540
102	cgccaagccg	ctgctgatca	tcgcccagga	cgtcgagggc	gaggcgctgt	ccaccctgg	600
104	cgtc						604
107	<210>	SEQ ID NO:	4				
108	<211>	LENGTH:	604				
109	<212>	TYPE:	DNA				
110	<213>	ORGANISM:	Mycobacterium aichiense				
112	<400>	SEQUENCE:	4				
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115	cgatgtcgcg	ggcgcacggca	ccaccaccgc	caccgtgctc	gctcaggcgc	tggtcgcga	120
117	aggtctgcgc	aacgtcgctg	ccggcgccaa	cccgcgtggc	ctgaagcgcg	gcacgagaa	180
119	ggccgtcgag	aagatcaccc	agacgctct	caagacgccc	aaggaggtcg	agaccaagga	240
121	ccagatcgcg	gccaccggcg	gatctccgc	gggcgaccag	accatcggt	acctgtatgc	300
123	cgaggccatg	gacaaggctcg	gcaacgaggg	tgtcatcacc	gtcgaggagt	ccaacacctt	360
125	cggcctgcag	ctcgagctca	ccgagggtat	gcccgtcgac	aagggttaca	tctcggttta	420
127	cttcgtgacc	gacgcccggc	gtcaggaagc	ggtcctcgag	gatccgtaca	tcctgctgg	480
129	gtcgtcgaaag	gttcgaccg	tcaaggacct	gcttccttg	ctggagaagg	tcatccagtc	540
131	ggccaagccg	ctgctgatca	tcgcccagga	cgtcgagggc	gaggccctgt	ccaccctgg	600
133	ggtc						604
136	<210>	SEQ ID NO:	5				
137	<211>	LENGTH:	604				
138	<212>	TYPE:	DNA				
139	<213>	ORGANISM:	Mycobacterium avium				
141	<400>	SEQUENCE:	5				
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144	cgacgtcgcc	ggtgacggca	cgacgacggc	cacggcgctc	gcccaggcgt	tggtcgcga	120
146	gggcctgcgc	aacgtcgccg	ccggcgccaa	cccgcgtggc	ctcaagcgcg	gcacgagaa	180
148	ggccgtcgag	aaggtcaccc	agaccctgct	caagacgccc	aaggaggtcg	agaccaagga	240
150	ccagatcgct	gccaccggcg	ccatctccgc	gggcgaccag	tcgatcgcc	acctgtatgc	300
152	cgaggccatg	gacaaggctcg	gcaacgaggg	cgtcatcacc	gtcgaggagt	ccaacacctt	360
154	cggcctgcag	ctcgagctca	ccgagggtat	gcccgtcgac	aagggttaca	tctcggttta	420
156	cttcgtcacc	gacgcccggc	gtcaggaagc	cgtcctcgag	gatccgtaca	tcctgctgg	480
158	cagctccaag	gttcgaccg	tcaaggacct	gctgccgctg	ctggagaagg	tcatccaggc	540

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160	cggcaagccg ctgctgatca tcgcccagga cgtcgaggc gaggccctgt ccaccctgg	600
162	cgtc	604
165	<210> SEQ ID NO: 6	
166	<211> LENGTH: 604	
167	<212> TYPE: DNA	
168	<213> ORGANISM: Mycobacterium bovis	
170	<400> SEQUENCE: 6	
171	ggaggatccg tacgagaaga tcggcgccga gctggtcaaa gaggtagcca agaagaccga	60
173	tgacgtcgcc ggtgacggca ccacgacggc caccgtctg gcccaggcgt tgggtcgca	120
175	gggcctgcgc aacgtcgccg ccggcgccaa cccgctcggt ctcaaacgcg gcataaaaa	180
177	ggccgtggag aaggtcaccg agaccctgt caagggcgcc aaggaggctg agaccaagga	240
179	gcagattgcg gccaccgcag cgatttcggc gggtgaccag tccatcggt acctgatcgc	300
181	cgaggcgatg gacaagggtgg gcaacgaggc cgtcatcacc gtcgaggagt ccaacacctt	360
183	tgggctgcag ctcgagctca ccgagggtat gcgggtcgac aagggttaca tctcgggta	420
185	tttcgtgacc gaccggagc gtcaggagc ggtcttggag gacccttaca tcctgctggt	480
187	cagctccaag gtgtccactg tcaaggatct gtcggcgctg ctcgagaagg tcatacgagc	540
189	cggtaagccg ctgctgatca tcgcccagga cgtcgaggc gaggcgctgt ccaccctgg	600
191	cgtc	604
194	<210> SEQ ID NO: 7	
195	<211> LENGTH: 604	
196	<212> TYPE: DNA	
197	<213> ORGANISM: Mycobacterium bovis BCG	
199	<400> SEQUENCE: 7	
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202	tgacgtcgcc ggtgacggca ccacgacggc caccgtctg gcccaggcgt tgggtcgca	120
204	gggcctgcgc aacgtcgccg ccggcgccaa cccgctcggt ctcaaacgcg gcataaaaa	180
206	ggccgtggag aaggtcaccg agaccctgt caagggcgcc aaggaggctg agaccaagga	240
208	gcagattgcg gccaccgcag cgatttcggc gggtgaccag tccatcggt acctgatcgc	300
210	cgaggcgatg gacaagggtgg gcaacgaggc cgtcatcacc gtcgaggagt ccaacacctt	360
212	tgggctgcag ctcgagctca ccgagggtat gcgggtcgac aagggttaca tctcgggta	420
214	tttcgtgacc gaccggagc gtcaggagc ggtcttggag gacccttaca tcctgctggt	480
216	cagctccaag gtgtccactg tcaaggatct gtcggcgctg ctcgagaagg tcatacgagc	540
218	cggtaagccg ctgctgatca tcgcccagga cgtcgaggc gaggcgctgt ccaccctgg	600
220	cgtc	604
223	<210> SEQ ID NO: 8	
224	<211> LENGTH: 604	
225	<212> TYPE: DNA	
226	<213> ORGANISM: Mycobacterium celatum Type 1	
228	<400> SEQUENCE: 8	
229	ggaggacccc tacaaaaaga tcggcgccga gctggtcaag gaagtcgcca agaagaccga	60
231	cgacgtcgcc ggtgacggta cgacgacggc cacgggtctg gcccaggcgt tggtaagga	120
233	gggcctgcgc aacgtcgccg ccggcgccaa cccgctcgcc ctgaagcgcc gcataaaaa	180
235	ggccgtcgag aaggtcaccg agacgctgt caagggcgcc aaggaggctg agaccaagga	240
237	gcagattgt gccaccgcgg ccatctccgc cggcgaccag tcgatcgcc acctgatcgc	300
239	cgaggccatg gacaaggctg gcaacgaggc cgtcatcacc gtcgaggagt ccaacacctt	360
241	cggcctgcag ctcgagctca ccgagggtat gcgggtcgac aagggttaca tctcgggta	420
243	tttcgtcacc gacgccccg gtcaggagc ggtgctcgag gagccgtaca tcctgctggt	480
245	cagctccaag gtgtcgacgg tcaaggactt gtcggcgctg ctggagaagg tcataccaggc	540
247	cggcaagccg ctgctgatca tcgcccagga cgtcgaggc gaaggccctct ccaccctgg	600

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Input Set : A:\280277 Sequence Listings.txt  
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249	cgtc	604
252	<210> SEQ ID NO: 9	
253	<211> LENGTH: 604	
254	<212> TYPE: DNA	
255	<213> ORGANISM: Mycobacterium celatum TypeII	
257	<400> SEQUENCE: 9	
258	ggaggacccc tacgagaaga tcggcgccga gctggtaag gaagtcgcca agaagaccga	60
260	cgacgtcgcg ggtgacggta cgacgacgac caccgtgtc gcccaggcg tggtaagaa	120
262	aggcctgcgc aacgtcgccg ccgggtccaa cccgctcgcc ctgaagcgcg gtatcgagaa	180
264	ggccgtcgag aaggtcaccg agacgctgtc caaggcgcc aaggaggtcg agaccaagga	240
266	gcagatcgct gccaccggg ccatctccgc cgggtgaccag tcgatcgccg acctgatcg	300
268	cgagggcgatg gacaaggctcg gcaacgaggg cgtcatcacc gtcgaggagt ccaacacctt	360
270	cgccctgcag ctcgagctca ccgagggtat gcgcttcgac aagggttaca tctcgggtta	420
272	cttcgtcacc gacgcccggc gtcaggaggc ggtgctcgag gagccctaca tcctgctgg	480
274	cagctccaag gtgtcgacgg tcaaggatct gctgccgtg ctggagaagg tcatccaggc	540
276	cggcaagccg ctgctgatca tcgcccggagga cgtcgagggt gaggcggtga gcaccctgg	600
278	cgtc	604
281	<210> SEQ ID NO: 10	
282	<211> LENGTH: 604	
283	<212> TYPE: DNA	
284	<213> ORGANISM: Mycobacterium chelonae	
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287	ggaggacccg tacgagaaga tcggcgctga gctggtaag gaagttgcca agaagactga	60
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291	aggctcgctg aacgtcgctg ccggcgccaa cccgctcgcc ctgaagcgcg gcacgagaa	180
293	ggccgtggag gccgtcacca gctctctgtc ggactccggc aaggagatcg acaccaagga	240
295	gcagatcgcg gccaccggg gcatctccgc ggggtgaccag tccatcggt atctgatcg	300
297	cgagggccatg gacaaggctcg gcaacgaggg tgtcatcacc gtcgaggagt ccaacacctt	360
299	cgccctgcag ctggagctca ccgagggtat gcgcttcgac aagggttaca tctcgggtta	420
301	cttcgtgacc gacgcccggc gtcaggaaac cgtctggag gatccctaca tcctgctgg	480
303	cagctccaag gtgtcgacgg tcaaggatct acttcccttg ctggagaagg tcatccaggc	540
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307	cgtc	604
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313	<213> ORGANISM: Mycobacterium chitae	
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320	aggctcgcc aacgtcgccg ccggcgccaa cccgctcgcc ctgaagcgcg gcacgagaa	180
322	ggccgtcgag accgtctcggt agaacctgtc caagtccggc aaggaggtcg agaccaagga	240
324	gcagatcgcc gccaccggc gatctccgc gggcgacacc accatcggt acctgatcg	300
326	cgagggccatg gacaagggtgg gcaacgaggg tgtcatcacc gtcgaggagt ccaacacctt	360
328	cgccctgcag ctggagctca ccgagggtat gcgcttcgac aagggttaca tctcgggtta	420
330	cttcgtgacc gacgcccggc gtcaggaaac cgtctggag gatccctaca tcctgctgg	480
332	cagctcgaaatctcgacccg tcaaggatct gctgccgtg ctggagaagg tcatccagtc	540
334	cggcaagccg ctgctgatca tcgcccggagga cgtcgagggc gaggccctgt cgaccctgg	600
336	ggtc	604

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339 <210> SEQ ID NO: 12  
 340 <211> LENGTH: 604  
 341 <212> TYPE: DNA  
 342 <213> ORGANISM: Mycobacterium microti  
 344 <400> SEQUENCE: 12

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347 tgacgtcgcc ggtgacggca ccacgacggc	caccgtgctg gcccaggcgt	tgggtcgca	120
349 gggcctgcgc aacgtcgccg cggcgccaa	cccgtcggt ctcaaacgcg	gcatcgaaaa	180
351 ggccgtggag aaggtcaccg agaccctgt	caagggcgcc aaggaggtcg	agaccaagga	240
353 gcagattgcg gccaccgcag cgatttcggc	gggtgaccag tccatcggtg	acctgatcgc	300
355 cgaggcgatg gacaagggtgg gcaacgaggg	cgtcatcacc gtcgaggagt	ccaacacctt	360
357 tgggctgcag ctcgagctca ccgagggtat	gcggttcgac aagggctaca	tctcgggta	420
359 ctctgtgacc gacccggagc gtcaggaggc	ggtcctggag gaccctata	tcctgcttgt	480
361 cagctccaag gtgtccactg tcaaggatct	gctgccgtg ctcgagaagg	tcatcgagc	540
363 cggtaagccg ctgctgatca tcgcccggagga	cgtcgagggc gaggcgctgt	ccacccttgt	600
365 cgtc			604
368 <210> SEQ ID NO: 13 369 <211> LENGTH: 604 370 <212> TYPE: DNA 371 <213> ORGANISM: Mycobacterium flavescens 373 <400> SEQUENCE: 13			
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376 cgacgtcgcc ggcgacggca ccaccaccgc	caccgtgctg gcccaggcgc	tcgtcgccga	120
378 gggctgcgc aacgtcgccg cggcgccaa	cccgtatggcg ctgaagcgcg	gtatcgagaa	180
380 ggccgtcgag aaggtcaccg agacgctgt	gaagtgcgc aaggaggtcg	agaccaagga	240
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386 cgggctgcag ctcgagctca ccgagggtat	gcggttcgac aagggctaca	tctcgggta	420
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390 gtcgtccaag gtgtcgaccg tcaaggatct	gctcccgttg ctggagaagg	tcattcaggc	540
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394 ggtc			604
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405 cgacgtcgcc ggcgacggca ccaccaccgc	caccgttctg gcacaggccc	tgggtcgta	120
407 aggtctgcgc aacgtcgctg cggcgccaa	cccgtcgcc ctgaagcgcg	gcatcgagaa	180
409 ggccgtcgag aaggtcaccg agacgctgt	gaagagcgcc aaggaggtgg	agaccaagga	240
411 gcagatcgct gccaccgcg gtatctccgc	cggtgaccag tccatcggtg	acctgatcgc	300
413 cgaggccatg gacaagggtcg gcaacgaggg	tgtcatcacc gtcgaggaga	gcaacacctt	360
415 cggcctgcag ctggagctca ccgggggtat	gcggttcgac aagggctaca	tctcgggcta	420
417 ctctgtgacc gacccggagc gtcaggaaagc	gatccctaca tcctgcttgt	480	
419 cagctccaag gtctcgaccg tcaaggact	gctgccgtg ctggagaagg	tcatccagtc	540
421 cggcaagccg ctgctgatca tcgcccggagga	cgtcgagggc gaaggccctgt	cgaccctggc	600
423 ggtc			604
426 <210> SEQ ID NO: 15			

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L:10 M:270 C: Current Application Number differs, Replaced Current Application No

L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date